



## CRF Problem Report

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 10516,361  
Filing Date: 11-30-2004  
Date Processed by STIC: 12-10-04

STIC Contact: Mark Spencer; Telephone: 571-272-2510; Fax: 571-273-0221

### Nature of Problem:

The CRF (was):

- Damaged or Unreadable (for Unreadable, see attached)  
 Blank (no files on CRF) (see attached)  
 Empty file (filename present, but no bytes in file) (see attached)  
 Virus-infected. Virus name: \_\_\_\_\_ The STIC will not process the CRF  
 Not saved in ASCII text  
 Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should only be the Sequence Listing.  
 Did not contain a Sequence Listing. (see attached sample)  
 Other.

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PLEASE USE THE CHECKER VERSION 4.2 PROGRAM TO REDUCE ERRORS.  
SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):  
U.S. Patent and Trademark Office, 220 20<sup>th</sup> Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

(Sample of submitted file)

10/5/16, 3618CT

SEQUENCE LISTING

1). GENERAL INFORMATION

i). NUMBER OF SEQUENCES : 31

2) INFORMATION FOR SEQ ID No.:1

i). SEQUENCE CHARACTERISTICS:

(A) LENGTH : 20 nucleotides

(A) TYPE : nucleic acid

(B) STANDARDNESS : single standard

(C) TOPOLOGY : linear

ii). MOLECULE TYPE : nucleic acid

iii). HYPOTHETICAL : yes

iv). ANTI-SENCE : no

(See sample Sequence Listing  
ON Next PAGE)

(Sample Sequence Listing)

<110> Smith, John; Smithgene Inc.  
<120> Example of a Sequence Listing  
<130> 01-00001  
<140> PCT/EP98/00001  
<141> 1998-12-31  
<150> US 08/999,999  
<151> 1997-10-15  
<160> 4  
<170> PatentIn version 2.0  
<210> 1  
<211> 389  
<212> DNA  
<213> Paramecium sp.  
<220>  
<221> CDS  
<222> (279)...(389)  
<300>  
<301> Doe, Richard  
<302> Isolation and Characterization of a Gene Encoding a  
Protease from Paramecium sp.  
<303> Journal of Genes  
<304> 1  
<305> 4  
<306> 1-7  
<307> 1988-06-31  
<308> 123456  
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tcatgtggca attgctggca gtgccacagg ctttcagcc aggcttaggg tgggttccgc 180  
cgccggcgccg cggccccctct cgcgctcctc tcgcgcctct ctctcgctct cctctcgctc 240

ggacctgatt	aggtagcag	gaggaggggg	cagttgc	atg	gtt	tca	atg	ttc	agc	29.6
				Met	Val	Ser	Met	Phe	Ser	
				1				5		

ttg	tct	ttc	aaa	tgg	cct	gga	ttt	tgt	ttg	ttt	gtt	tgt	ttg	ttc	caa	344
Leu	Ser	Phe	Lys	Trp	Pro	Gly	Phe	Cys	Leu	Phe	Val	Cys	Leu	Phe	Gln	
			10					15					20			

tgt	ccc	aaa	gtc	ctc	ccc	tgt	cac	tca	tca	ctg	cag	ccg	aat	ctt	389
Cys	Pro	Lys	Val	Leu	Pro	Cys	His	Ser	Ser	Leu	Gln	Pro	Asn	Leu	
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<212> PRT  
<213> Paramecium sp.

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1				5					10					15		
Phe	Val	Cys	Leu	Phc	Gln	Cys	Pro	Lys	Val	Leu	Pro	Cys	His	Ser	Ser	
		20						25					30			
Leu	Gln	Pro	Asn	Leu												
		35														

<210> 3  
<211> 11  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Designed peptide based on size and polarity to act as a linker between the alpha and beta chains of Protein XYZ.

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1				5												

<210> 4  
<400> 4  
000

[Annex VIII follows]